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          GenCore version 4.5

          Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

Run on:    December 26, 2001, 10:33:29 ; Search time 15.89 Seconds
                                          (without alignments)
                                          662.778 Million cell updates/sec

Title:              US-09-497-967-7
Perfect score:      2540
Sequence:           1 MKKNILVILIISLFINQIKS.........QCDFANFLSISLLLISYYLL 468

Scoring table:      BLOSUM62
                    Gapop 10.0 , Gapext 0.5

Searched:           212252 seqs, 22503292 residues

Total number of hits satisfying chosen parameters:    212252

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :    Issued_Patents_AA:*
        1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
        2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
        3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
        4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
        5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
        6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:**

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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SUMMARIES

Result No.	Score	Query Match %	Length	DB	ID	Description
1	200.5	7.9	1345	2	US-08-977-767-3	Sequence 3, Appli
2	187.5	7.4	801	1	US-07-906-349A-6	Sequence 6, Appli
3	179.5	7.1	341	2	US-08-209-521-11	Sequence 11, Appl
4	175.5	6.9	3075	2	US-08-460-309-5	Sequence 5, Appli
5	175.5	6.9	3075	2	US-08-125-077-5	Sequence 5, Appli
6	166.5	6.6	969	2	US-08-284-941-2	Sequence 2, Appli
7	166.5	6.6	969	2	US-08-447-642-2	Sequence 2, Appli
8	166.5	6.6	969	4	US-09-236-503-2	Sequence 2, Appli
9	166.5	6.6	969	5	PCT-US93-02147A-2	Sequence 2, Appli
10	157	6.2	3111	2	US-08-460-309-4	Sequence 4, Appli
11	157	6.2	3111	2	US-08-125-077-4	Sequence 4, Appli
12	150.5	5.9	1111	4	US-08-317-450B-15	Sequence 15, Appl
13	150.5	5.9	1111	1	US-08-800-593-15	Sequence 15, Appl
14	150.5	5.9	1193	4	US-08-317-450B-13	Sequence 13, Appl
15	150.5	5.9	1193	1	US-08-800-593-13	Sequence 13, Appl
16	146	5.7	355	1	US-08-292-549-6	Sequence 6, Appli
17	146	5.7	355	4	US-09-006-353A-14	Sequence 14, Appli
18	143.5	5.6	610	2	US-09-365-470-3	Sequence 3, Appli
19	143.5	5.6	610	3	US-09-209-668-19	Sequence 19, Appl
20	143.5	5.6	610	4	US-09-009-490A-89	Sequence 89, Appl
21	141.5	5.6	610	6	5217870-2	Patent No. 5217870
22	138	5.4	2471	1	US-08-185-432-16	Sequence 16, Appl
23	138	5.4	2471	1	US-08-083-590A-19	Sequence 19, Appli
24	138	5.4	2471	3	US-08-532-384-19	Sequence 19, Appl
25	135.5	5.3	2523	1	US-08-185-432-18	Sequence 18, Appli
26	132	5.2	1147	1	US-08-144-121-3	Sequence 3, Appli
27	132	5.2	1147	2	US-08-735-893-3	Sequence 3, Appli
28	131.5	5.2	314	2	US-08-460-309-19	Sequence 19, Appl
29	131.5	5.2	314	2	US-08-125-077-19	Sequence 19, Appl
30	131.5	5.2	1466	6	5256642-6	Patent No. 5256642
31	131.5	5.2	1466	6	5472939-6	Patent No. 5472939
32	131.5	5.2	1537	6	5256642-5	Patent No. 5256642
33	131.5	5.2	1537	6	5472939-5	Patent No. 5472939
34	131.5	5.2	1847	6	5256642-10	Patent No. 5256642
35	131.5	5.2	1847	6	5472939-10	Patent No. 5472939
36	131.5	5.2	2039	6	5256642-2	Patent No. 5256642
37	131.5	5.2	2039	6	5472939-2	Patent No. 5472939
38	130.5	5.1	581	3	US-08-665-259-23	Sequence 23, Appl
39	130.5	5.1	581	2	US-08-762-500-23	Sequence 23, Appl
40	130	5.1	830	6	5378464-3	Patent No. 5378464
41	129	5.1	574	6	5378464-3	Patent No. 5378464
42	128.5	5.1	830	1	US-08-110-158-4	Sequence 4, Appli
43	127.5	5.0	581	3	US-08-152-019A-27	Sequence 27, Appl
44	127.5	5.0	581	1	US-08-482-677-6	Sequence 6, Appli
45	127.5	5.0	581	3	US-08-650-599A-2	Sequence 2, Appli

ALIGNMENTS

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RESULT 1
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; Sequence 3, Application US/08977767
; Patent No. 5972684
; GENERAL INFORMATION:
;   APPLICANT: Bandman, Olga
;   APPLICANT: Yue, Henry
;   APPLICANT: Greenwald, Sara
;   APPLICANT: Corley, Neil C.
;   TITLE OF INVENTION: CARBONIC ANHYDRASE VIII
;   NUMBER OF SEQUENCES: 3
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: Incyte Pharmaceuticals, Inc.
;     STREET: 3174 Porter Drive
;     CITY: Palo Alto
;     STATE: CA
;     COUNTRY: USA
;     ZIP: 94304
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE: Diskette
;     COMPUTER: IBM Compatible
;     OPERATING SYSTEM: DOS
;     SOFTWARE: FastSEQ for Windows Version 2.0
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER: US/08/977,767
;     FILING DATE: Herewith
;     CLASSIFICATION: 424
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:
;     FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;     NAME: Billings, Lucy J.
;     REGISTRATION NUMBER: 36,749
;     REFERENCE/DOCKET NUMBER: PF-0423 US
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: 650-855-0555
;     TELEFAX: 650-845-4166
;     TELEX:
; INFORMATION FOR SEQ ID NO: 3:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 1345 amino acids
;     TYPE: amino acid
;     STRANDEDNESS: single
;     TOPOLOGY: linear
;   IMMEDIATE SOURCE:
;     LIBRARY: GenBank
;     CLONE: 1532042
US-08-977-767-3
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Query Match    7.9%; Score 200.5; DB 2; Length 1345;
Best Local Similarity  25.8%; Pred. No. 3.3e-09;
Matches 111; Conservative 5; Mismatches 212; Indels 103; Gaps 17;

Qy   21 ANCPVGTETNTAGQVDDLG-TPANCVNCQKNFYYNNAAAFVPGAS---TCTPCPQKKDAG 76
Db  649 AGCACGGACGAGCGGCTGGCTCTGAACACCAGATCAACCAGATCAACCATCAGGTTTCTCTGCTGAGGTG 708

Qy   77 AQPNPPATANLVTQCNVKCPAG----TAIAGGATDYAAIITECVNCRINFYNENAPNFN 131
Db  709 CAACTCATCCACTTCAACCAAGAACTCTATGGGAACCTCAGTGCCGCCA-------CC 759

Qy  132 AGASTCTACPVNRVGGALTAGNAAT-IVAQCNVACPFGTALDDGVTTDYVRSFTECVKC 189
Db  760 AGGGGCCCCAA--TGGCCTGGCCATTCTCAGCCTCTTTGTCAATGT--------- 803

Qy  190 RLNFYYNGNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAA 249
Db  804 ------GGCTG------GTAGCT-CAAACCGTTCCTCAGACGCCTCCTTAACCGTGACA 850

Qy  250 GVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCN 309
Db  851 C---------CATCA---------CCCGCATCTCC------TATAAGAATGATGCC 882

Qy  310 IACPDGTAIASGATNYVIILQTECLNCAANFYFDGNNFQAGSSRCKACPANKV----QGAV 365
Db  883 TAC------------TTTCTTCAAGACCTGAGCCTGGAGCTCGTGTGCCCGAGTC 926

Qy  366 ATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFYTTKQTDWVAGIDTC 425
Db  927 CTTTGGCTTCATCA--CCTATCAGGGCTCTCTCAGCAGCCTCGTCTCTGCT--CGGAGAC 980

Qy  426 TSCNKKLTSGA 436
Db  981 TGTTACCTGGA 991
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RESULT 2
US-07-906-349A-6
; Sequence 6, Application US/07906349A
; Patent No. 5434064
; GENERAL INFORMATION:
; APPLICANT: Schlessinger, Joseph
; APPLICANT: Skolnik, Edward Y.
; APPLICANT: Margolis, Benjamin L.
; TITLE OF INVENTION: A NOVEL EXPRESSION-CLONING METHOD FOR
; TITLE OF INVENTION: IDENTIFYING TARGET PROTEINS FOR EUKARYOTIC TYROSINE KINASES AN
; TITLE OF INVENTION: TARGET PROTEINS
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Browdy and Neimark
; STREET: 419 Seventh Street, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20004
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/906,349A
; FILING DATE: 30-JUN-1992
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/643,237
; FILING DATE: 18-JAN-1991
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-628-5197
; TELEFAX: 202-737-3528
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:

Right column

; LENGTH: 801 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-07-906-349A-6

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Query Match    7.4%; Score 187.5; DB 1; Length 801;
Best Local Similarity  25.7%; Pred. No. 2.3e-08;
Matches 116; Conservative 7; Mismatches 195; Indels 133; Gaps 21;

Qy   21 ANCPVGT-----ETNTAGQV-DDLGTTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKK 73
Db  253 AGCCTGTGAGGGAGTGCAGAAACACCGTTTC------------TTAATTGGGTGTTCGACG 302

Qy   74 DAGAQPNPPATAN--LVTQCNVKCPAGTAIAG-GATDYAAIITECVNCRINFYNENAPNF 130
Db  303 GAGACTGTCGTCCGACACTCCCTCACGTCTTGTGGTGCGCGGGTCACCCT------- 352

Qy  131 NAGASTCT-ACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECVKC 189
Db  353 GTGACCCTCTCACTTGG--TTGGA------------ACTT-TAGGGG------- 385

Qy  190 RLNFYYNGNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDA---TITAQCNVACPDGTI 246
Db  386 ------GTGGGAGGGGCACCGGCCCAGTGGACACTGGGAGTGAAACCAAC--CTT 436

Qy  247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGGAAT--- 303
Db  437 GAAA-------TCCCCCA--------CCCTCCCCCGGTTGGATTAAAAATGCC 475

Qy  304 LAKQCNIACPDGTAIASGA------TNYVILQTECLNCAANFYFDGNNFQAG 349
Db  476 AAAACTTACCTATAAATTAAGAAGAGTTTTTATTACAAATTCAACCT------AAA 526

Qy  350 SSRCKACPANKVQGAVATAGGTATLIA----QCALECPAGTVLTDGTTSTYKQAASECV 404
Db  527 TTTTTACGGTTTTGA-ATGGATATTTAATTCTTCTCAAAAATAATGTTTAAATCACTGCT 585

Qy  405 KCAANFYTTKQTDWVAGIDTCTSCNKKLTSG 435
Db  586 GCTCCTCTT--------TCCCCTCCTTTG 606
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RESULT 3
US-08-209-521-11
; Sequence 11, Application US/08209521
; Patent No. 5922855
; GENERAL INFORMATION:
; APPLICANT: Liskay, Robert M.
; APPLICANT: Bronner, C. Eric
; APPLICANT: Baker, Sean M.
; APPLICANT: Bollag, Roni J.
; APPLICANT: Kolodner, Richard D.
; TITLE OF INVENTION: MAMMALIAN DNA MISMATCH REPAIR GENES
; TITLE OF INVENTION: hMLH1 AND hPMS1
; NUMBER OF SEQUENCES: 30
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Kolisch, Hartwell, Dickinson, McCormack &
; ADDRESSEE: Heuser
; STREET: 520 S.W. Yamhill, Suite 200
; CITY: Portland
; STATE: Oregon
; COUNTRY: US
; ZIP: 97204
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/209,521

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; FILING DATE: 08-MAR-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
;   NAME: Van Rysselberghe, Pierre C.
;   REGISTRATION NUMBER: 33,557
;   REFERENCE/DOCKET NUMBER: OHSU 306A
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (503) 224-6655
;   TELEFAX: (503) 295-6679
; INFORMATION FOR SEQ ID NO: 11:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 341 amino acids
;     TYPE: amino acid
;     STRANDEDNESS: single
;     TOPOLOGY: linear
;   MOLECULE TYPE: cDNA
;   POSITION IN GENOME:
;     MAP POSITION: 3p21.3-23
US-08-209-521-11
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Query Match         7.1%;  Score 179.5; DB 2; Length 341;
Best Local Similarity   25.8%; Pred. No. 3.8e-08;
Matches  98; Conservative   9; Mismatches 130; Indels 143; Gaps  20;

Qy   134 ASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECVKCRLNF 193
             |||| :||  |  ||  |||  ||  | |
Db     3 ATTCAAAG---AGATTTGGAAAATGAGTAACAT------GATTATTTACT-CATC---- 47

Qy   194 YYNGNGNGTPFNPGKSQCT-PCPAIKPANVAQATLGNDATI-----------TAQCNV 239
          |  : |   ||  : |       ||        ||    |
Db    48 ------TTTTTGGTATCTAACAAAAGAAGAGATCTGGATATTGTATGTGAAAAGGKTCAC-T 99

Qy   240 ACPDGTISAAGVNNWVAQNFECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAG 299
          || |  |||  :  ||  |  ||  |       |     ||   |   |
Db   100 ACTAGTAAA--------------CTGCA----------GTC--CTTTGAGGADTTTA 130

Qy   300 G--AATLAKQCNIAC------------------PDGTA----IASGATNYVILQTECLNCAA 337
          |  |  |     | |                  || |       |||
Db   131 GCCAGTATTTCTACCTATGGCTTTCGAGGTGAGGTAAGCTAAAGAT--------TCAA 180

Qy   338 NEYFDGNNFQAGSSR--CKACPANKVQGAVATAG---------------------GTATLIAQ 377
          |||                :  ||  ||||
Db   181 GAAATGTKTAAAATATCCTCCTGTGATGACATTGTYTGTCATTTGTTGTAGTAGTTATTTCT 240

Qy   378 CALECPAG------------------------TVLTDGTTSTYKQAASECVKCAANFYTTKQTDW 418
          |                                | :   ||           ||
Db   241 CAACATAGATAAATAAGGTTTGGTACCTTTTACTTGTTAAATGTATGCAA--ATGTATGC 298

Qy   419 VAGIDTCT--SCNKKLTSGA 436
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Db   299 AAA--TCTGHGCAAACTTAA 316
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;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/460,309
;   FILING DATE:
;   CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US 08/125,077
;   FILING DATE: 22-SEP-1993
;   APPLICATION NUMBER: US PCT/US 94/10730
;   FILING DATE: 21-SEP-1994
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US 07/472,319
;   FILING DATE: 30-JAN-1990
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US 07/919,951
;   FILING DATE: 27-JUL-1992
; ATTORNEY/AGENT INFORMATION:
;   NAME: Campbell, Cathryn A.
;   REGISTRATION NUMBER: 31,815
;   REFERENCE/DOCKET NUMBER: P-LA 9721
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (619) 535-9001
;   TELEFAX: (619) 535-8949
; INFORMATION FOR SEQ ID NO: 5:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 3075 amino acids
;     TYPE: amino acid
;     TOPOLOGY: linear
US-08-460-309-5
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Best Local Similarity   21.2%; Pred. No. 1.6e-06;
Matches 110; Conservative   44; Mismatches 187; Indels 177; Gaps  30;

Qy    18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA 77
          :   |         :|   :   || :         :: || |       ||
Db   706 VEHCECPQGY--------------TGTSCESCLSGYYRVDGILF---GGICQPCECHGHA-- 748

Qy    78 QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC 137
          :::||         |||      || | :| || ||  ||| ||
Db   749 ------------AECNVH---GVCI---ACAHNTTGVHCEQCLPGFYGEPSRGTPGDCQPC 791

Qy   138 TACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDG--VTTDYVR--SFTECVKCRLN 192
          |||         ::        :    :||:  |  ||    :|:
Db   792 -ACPL----------TIAS--NNFSPT-CHLNDGDEVVCDWCAPGYSGAWCERCADG 834

Qy   193 FYYNGNGNGTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVN 252
          :                  |    ::|      |  :: :: |  |  |
Db   835 YYGN-------PTVPGES-CVPCDC--SGNVDPSEAGHCDSVTGEC-LKCLGNTDGA---- 880

Qy   253 NWVAQNTECTNCAPNFYNN-NAPN--------------------FNPGNSTCLPCPANKD-- 291
          |              :|:  | |                     |
Db   881 ------HCERCADGFYGDAVTAKNCRACECHVKGSHSAVCHLETGLCDCKPNVTGQQCD 933

Qy   292 -------YGAEATAGGAATLAKQCNIACPDGTAIASGATNY-----VILQTECLNCAAN 338
          |                    :: :: | |         |
Db   934 QCLHGYYGLDSGHG-----CRPCN--CSVAGSVSDGCTDEGQCHCVPGVAGKRCDRCAHG 986

Qy   339 FYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCALECPAGT------VLTDGT 392
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Db   987 FY----AYQDGS--CTPCDCPHTQNTCDPETG------ECV--CPPHTQGVKCEECEDGH 1032

Qy   393 TSTYKQAASECVKCAANFYTTKQTDWVAGIDTCTS------------------C 428
          |      :: |           |  |   |:  |:
Db  1033 WGYDAEVGCQACNCSLVGSTHHRCDVVTGHCQCKSKFGGRACVQCSLGYRDFPDCVPCDC 1092

Qy   429 NKKLTSGAFANLPES---------------AKKNI---QCD 451
          | ||||  || ::||                :||   ||
Db  1093 DLRGTSGDACNLEQGLCGCVEETGACPCKENVFGPQCN 1130
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RESULT   4
US-08-460-309-5
; Sequence 5, Application US/08460309
; Patent No. 5837496
; GENERAL INFORMATION:
;   APPLICANT: Engvall, Eva
;   APPLICANT: Leivo, Ilmo
; TITLE OF INVENTION: Nucleic Acids Encoding Merosin, Merosin
; TITLE OF INVENTION: Fragments and Uses Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Campbell and Flores
;   STREET: 4370 La Jolla Village Drive, Suite 700
;   CITY: San Diego
;   STATE: California
;   COUNTRY: USA
;   ZIP: 92122
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
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us-09-497-967-7.rai

RESULT 5
US-08-125-077-5
; Sequence 5, Application US/08125077
; Patent No. 5872231
; Patent No. 5872231 5840863
; GENERAL INFORMATION:
; APPLICANT: Engvall, Eva
; APPLICANT: Leivo, Ilmo
; TITLE OF INVENTION: Nucleic Acids Encoding Merosin, Merosin
; TITLE OF INVENTION: Fragments and Uses Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Campbell and Flores
;   STREET: 4370 La Jolla Village Drive, Suite 700
;   CITY: San Diego
;   STATE: California
;   COUNTRY: USA
;   ZIP: 92122
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/125,077
;   FILING DATE: 22-SEP-1993
;   CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US PCT/US 94/10730
;   FILING DATE: 21-SEP-1994
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US 07/472,319
;   FILING DATE: 30-JAN-1990
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US 07/919,951
;   FILING DATE: 27-JUL-1992
; ATTORNEY/AGENT INFORMATION:
;   NAME: Campbell, Cathryn A.
;   REGISTRATION NUMBER: 31,815
;   REFERENCE/DOCKET NUMBER: P-LA 9721
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (619) 535-9001
;   TELEFAX: (619) 535-8949
; INFORMATION FOR SEQ ID NO: 5:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 3075 amino acids
;   TYPE: amino acid
;   TOPOLOGY: linear
US-08-125-077-5

Query Match   6.9%; Score 175.5; DB 2; Length 3075;
Best Local Similarity 21.2%; Pred. No. 1.6e-06;
Matches 110; Conservative 44; Mismatches 187; Indels 177; Gaps 30;

Qy   18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA 77
             ::|| | |:   :| :   :| : | | || :||: :
Db   706 VEHCECPQGY----------TGTSCESCLSGYYRVDGILF--GGICQPCECHGHA-- 748

Qy   78 QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC 137
             :||| ||| |: | ||||| | : || |::|| ::| | ||
Db   749 --------------AECNVH---GVCI---ACAHNTGVHCEQCLPGFYGEPSRGTPGDCQPC 791

Qy   138 TACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDG--VTTDYVR---SFTECVKCRLN 192
             || :   :|| | || || || || :   :: |:| :| |:|:
Db   792 -ACPL----------TIAS--NNFSPT-CHLNDGDEVVCDWCAPGYSGAWCERCADG 834

Qy   193 FYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVN 252
             |:||:|||| |: | | : | | :   ::| :| | :|
Db   835 YYGN-------PTVPGES-CVPCDC--SGNVDPSEAGHCDSVTGEC-LKCLGNTDGA---- 880

Qy   253 NWVAQNTECTNCAPNFYNN-NAPN-------------FNPGNSTCLPCPANKD------- 291
             : ||
Db   881 ------HCERCADGFYGDAVTAKNCRACECHVKGSHSAVCHLEIGLCDCKPNVTGQQCD 933

Qy   292 -----YGAEATAGGAATLAKQCNIACPDGTAIASGATNY------VILQTECLNCAAN 338
             ::   ::| :| || | : ||:|:::| :: :||
Db   934 QCLHGYYGLDSGHG-----CRPCN--CSVAGSVSDGCTDEGQCHCVPGVAGKRCDRCAHG 986

Qy   339 FYFDGNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCALECPAGT------VLTDGT 392
             | ||| : || |: : | | |: || : | | | | :| ||
Db   987 FY---AYQDGS--CTPCDCPHTQNTCDPETG-----ECV-CPPHTQGVKCEECEDGH 1032

Qy   393 TSTYKQAASECVKCAANFYTTKQTDWVAGIDTCTS---------------------C 428
             |: :| ||| | |::   :   : : |:| |
Db   1033 WGYDAEVGCQACNCSLVGSTHHRCDVVTGHCQCKSKFGGRACVQCSLGYRDFPDCVPCDC 1092

Qy   429 NKKLTSGAEANLPES-----------AKKNI---QCD 451
             : ::||| || |: |||
Db   1093 DLRGTSGDACNLEQGLCGCVEETGACPCKENVFGPQCN 1130

RESULT 6
US-08-284-941-2
; Sequence 2, Application US/08284941
; Patent No. 5863756
; GENERAL INFORMATION:
; APPLICANT: BARR, PHILIP J
; APPLICANT: KIEFER, MICHAEL C
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PACE 4 AND
; TITLE OF INVENTION: PACE 4.1 GENE AND POLYPEPTIDES IN CELLS
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
;   ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM
;   STREET: FIVE PALO ALTO SQUARE
;   CITY: PALO ALTO
;   STATE: CALIFORNIA
;   COUNTRY: USA
;   ZIP: 94306
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/284,941
;   FILING DATE: 2 August 1994
;   CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
;   NAME: NEELEY PH.D., RICHARD L.
;   REGISTRATION NUMBER: 30092
;   REFERENCE/DOCKET NUMBER: CHIR-009/01US
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (415) 843-5070
;   TELEFAX: (415) 857-0663
;   TELEX: 380816 COOLEY PA
; INFORMATION FOR SEQ ID NO: 2:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 969 amino acids
;   TYPE: amino acid
;   TOPOLOGY: linear
;   MOLECULE TYPE: protein
US-08-284-941-2

Query Match   6.6%; Score 166.5; DB 2; Length 969;
Best Local Similarity 20.7%; Pred. No. 2.1e-06;
Matches 86; Conservative 53; Mismatches 135; Indels 141; Gaps 25;

Qy   70 PQKKDAGAQPNPPATANLV-TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
           |: :: | :|| : ||:: | :   : |:   ::|| ::|
Db   672 PEDEEDYTAQSTPGSANILQTSVCHPEC-GDKGCDGPNAD-----QCLNC-VHF--SL 720

Qy   128 PNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV 187
           : || || | | | | || | | | ::

Db 721 GSVKTSRKCVSVCPLGYFG-----DTAARRCRCHKGCETCSS--------RAATQCL 765

Qy 188 KCRLNFYYNGNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
Db 766 SCRRGFYIH-QEMNT------CVTLCPAGFYADESQKN------CLKCHPSC----- 804

Qy 247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGAEATAGGAATLA 305
Db 805 -----KKCVDEPEKCTVC------KEGFSLARGSCIPDCEPGTYFDSELIRCG------ 846

Qy 306 KQCNIACPDGTAIASGATNYVILQTECLNCAANFYD------GNNFQAGS-------S 351
Db 847 -ECHHTC--GTCVGPG------REECIHCAKNFHFHDWKCVPACCGEGFYPEEMPGLPHK 896

Qy 352 RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY 411
Db 897 VCRRCDEN-----------------CLSC-AG------SSRNCSRCKTGF- 922

Qy 412 TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD--FANFLSISLLL 462
Db 923 ------TQLGTSCITNHTCSNADETFCEMVKSNRLCERKLFIQFCCRTCLL 967

RESULT 7
US-08-447-642-2
; Sequence 2, Application US/08447642
; Patent No. 5989890
; GENERAL INFORMATION:
; APPLICANT: BARR, PHILIP J
; APPLICANT: KIEFER, MICHAEL C
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PACE 4 AND
; TITLE OF INVENTION: PACE 4.1 GENE AND POLYPEPTIDES IN CELLS
; NUMBER OF SEQUENCES: 16
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM
; STREET: FIVE PALO ALTO SQUARE
; CITY: PALO ALTO
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94306
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/447,642
; FILING DATE: 23-MAY-1995
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/284,941
; FILING DATE: 2 August 1994
; ATTORNEY/AGENT INFORMATION:
; NAME: NEELEY PH.D., RICHARD L.
; REGISTRATION NUMBER: 30092
; REFERENCE/DOCKET NUMBER: CHIR-009/01US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 843-5070
; TELEFAX: (415) 857-0663
; TELEX: 380816 COOLEY PA
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 969 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-447-642-2

Query Match   6.6%; Score 166.5; DB 2; Length 969;
Best Local Similarity   20.7%; Pred. No. 2.1e-06;
Matches   86; Conservative   86; Mismatches 53; Mismatches 135; Indels 141; Gaps 25;

Qy 70 PQKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
Db 672 PEDEEDYTAQSTPGSANIIQTSVCHPEC-GDKGCDGPNAD-----QCLNC-VHF---SL 720

Qy 128 PNFNAGASTCACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV 187
Db 721 GSVKTSRKCVSVCPLGYFG-----DTAARRCRCHKGCETCSS--------RAATQCL 765

Qy 188 KCRLNFYYNGNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
Db 766 SCRRGFYIH-QEMNT------CVTLCPAGFYADESQKN------CLKCHPSC----- 804

Qy 247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGAEATAGGAATLA 305

Qy 70 PQKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
Db 672 PEDEEDYTAQSTPGSANIIQTSVCHPEC-GDKGCDGPNAD-----QCLNC-VHF---SL 720

Qy 128 PNFNAGASTCACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV 187
Db 721 GSVKTSRKCVSVCPLGYFG-----DTAARRCRCHKGCETCSS--------RAATQCL 765

Qy 188 KCRLNFYYNGNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
Db 766 SCRRGFYIH-QEMNT------CVTLCPAGFYADESQKN------CLKCHPSC----- 804

Qy 247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGAEATAGGAATLA 305
Db 805 -----KKCVDEPEKCTVC------KEGFSLARGSCIPDCEPGTYFDSELIRCG------ 846

Qy 306 KQCNIACPDGTAIASGATNYVILQTECLNCAANFYD------GNNFQAGS-------S 351
Db 847 -ECHHTC--GTCVGPG------REECIHCAKNFHFHDWKCVPACCGEGFYPEEMPGLPHK 896

Qy 352 RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY 411
Db 897 VCRRCDEN-----------------CLSC-AG------SSRNCSRCKTGF- 922

Qy 412 TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD--FANFLSISLLL 462
Db 923 ------TQLGTSCITNHTCSNADETFCEMVKSNRLCERKLFIQFCCRTCLL 967

RESULT 8
US-09-236-503-2
; Sequence 2, Application US/09236503
; Patent No. 6277590
; GENERAL INFORMATION:
; APPLICANT: Barr, Philip J
; APPLICANT: Kiefer, Michael C
; TITLE OF INVENTION: Compositions and Methods for PACE 4 and 4.1 Gene and
; TITLE OF INVENTION: Polypeptides in Cells
; FILE REFERENCE: CHIR-009/04US
; CURRENT APPLICATION NUMBER: US/09/236,503
; CURRENT FILING DATE: 1999-01-25
; EARLIER APPLICATION NUMBER: 08/447,642
; EARLIER FILING DATE: 1995-05-23
; EARLIER APPLICATION NUMBER: 08/284,941
; EARLIER FILING DATE: 1994-08-02
; EARLIER APPLICATION NUMBER: 07/848,629
; EARLIER FILING DATE: 1992-03-09
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
; LENGTH: 969
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-236-503-2

Query Match   6.6%; Score 166.5; DB 4; Length 969;
Best Local Similarity   20.7%; Pred. No. 2.1e-06;
Matches   86; Conservative   53; Mismatches 135; Indels 141; Gaps 25;

Qy 70 PQKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA 127
Db 672 PEDEEDYTAQSTPGSANIIQTSVCHPEC-GDKGCDGPNAD-----QCLNC-VHF---SL 720

Qy 128 PNFNAGASTCACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV 187
Db 721 GSVKTSRKCVSVCPLGYFG-----DTAARRCRCHKGCETCSS--------RAATQCL 765

Qy 188 KCRLNFYYNGNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI 246
Db 766 SCRRGFYIH-QEMNT------CVTLCPAGFYADESQKN------CLKCHPSC------ 804

Qy 247 SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGAEATAGGAATLA 305

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Qy  188  KCRLNFYYNGNGNTPFNPGKSQC-TPCPAIKPANVAQATLGNDATITAQCNVACPDGTI  246
              ||  ||::|       ||::|        |:   ||  |   || : :|     |:|  ||||  |:: :||
Db  766  SCRRGFYHH-QEMNT------------------CVTLCPAGFYADESQKN--------CLKCHPSC-  804

Qy  247  SAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLP-CPANKDYGAEATAGGAATLA  305
                        ||  |  :|:|||        |  :||
Db  805  ----KKCVDEPEKCTVC--------KEGFSLARGSCIPDCEPGTYFDSELIRCG-------  846

Qy  306  KQCNIACPDGTAIASGATNYVILQTECLNCAANFYD--------GNNFQAGS------S  351
              |||::||||     |  ||||||:||||||     ||:|||||:|||     |||||||:||:||
Db  847  -ECHHTC--GTCVGPG------REECIHCAKNFHFHDWKCVPACGEGFYPEEMPGLPHK  896

Qy  352  RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY  411
              |:| ||                 ||      ||  |   |||  ||:||
Db  897  VCRRCDEN--------------------------------CLSC-AG----------SSRNCSRCKTGF-  922

Qy  412  TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD---FANFLSISLLL  462
              ||||||||||||||||||||||||:||       ||  |||: :| || : ||:||
Db  923  -------TQLGTSCITNHTCSNADETFCEMVKSNRLCERKLFIQFCCRTCLL  967


RESULT  10
US-08-460-309-4
;  Sequence 4, Application US/08460309
;  Patent No. 5837496
;  GENERAL INFORMATION:
;      APPLICANT: Engvall, Eva
;      APPLICANT: Leivo, Ilmo
;      TITLE OF INVENTION: Nucleic Acids Encoding Merosin, Merosin
;      TITLE OF INVENTION: Fragments and Uses Thereof
;      NUMBER OF SEQUENCES: 23
;      CORRESPONDENCE ADDRESS:
;          ADDRESSEE: Campbell and Flores
;          STREET: 4370 La Jolla Village Drive, Suite 700
;          CITY: San Diego
;          STATE: California
;          COUNTRY: USA
;          ZIP: 92122
;      COMPUTER READABLE FORM:
;          MEDIUM TYPE: Floppy disk
;          COMPUTER: IBM PC compatible
;          OPERATING SYSTEM: PC-DOS/MS-DOS
;          SOFTWARE: PatentIn Release #1.0, Version #1.25
;      CURRENT APPLICATION DATA:
;          APPLICATION NUMBER: US/08/460,309
;          FILING DATE:
;          CLASSIFICATION: 435
;      PRIOR APPLICATION DATA:
;          APPLICATION NUMBER: US 08/125,077
;          FILING DATE: 22-SEP-1993
;          APPLICATION NUMBER: US PCT/US 94/10730
;          FILING DATE: 21-SEP-1994
;      PRIOR APPLICATION DATA:
;          APPLICATION NUMBER: US 07/472,319
;          FILING DATE: 30-JAN-1990
;      PRIOR APPLICATION DATA:
;          APPLICATION NUMBER: US 07/919,951
;          FILING DATE: 27-JUL-1992
;      ATTORNEY/AGENT INFORMATION:
;          NAME: Campbell, Cathryn A.
;          REGISTRATION NUMBER: 31,815
;          REFERENCE/DOCKET NUMBER: P-LA 9721
;      TELECOMMUNICATION INFORMATION:
;          TELEPHONE: (619) 535-9001
;          TELEFAX: (619) 535-8949
;  INFORMATION FOR SEQ ID NO: 4:
;      SEQUENCE CHARACTERISTICS:
;          LENGTH: 3111 amino acids
;          TYPE: amino acid
;          TOPOLOGY: linear
;  US-08-460-309-4
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Db  805  ----KKCVDEPEKCTVC--------KEGFSLARGSCIPDCEPGTYFDSELIRCG-----  846
                        ||  |  :|:|||
Qy  306  KQCNIACPDGTAIASGATNYVILQTECLNCAANFYD--------GNNFQAGS------S  351
              |||::||||     |  ||||||:||||||     ||:|||||:|||
Db  847  -ECHHTC--GTCVGPG------REECIHCAKNFHFHDWKCVPACGEGFYPEEMPGLPHK  896

Qy  352  RCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFY  411
              |:| ||
Db  897  VCRRCDEN--------------------------------CLSC-AG----------SSRNCSRCKTGF-  922

Qy  412  TTKQTDWVAGIDTCTSCNKKLT-SGAEANLPESAKKNIQCD---FANFLSISLLL  462
              ||||||||||||||||||||||||:||
Db  923  -------TQLGTSCITNHTCSNADETFCEMVKSNRLCERKLFIQFCCRTCLL  967


RESULT  9
PCT-US93-02147A-2
;  Sequence 2, Application PC/TUS9302147A
;  GENERAL INFORMATION:
;      APPLICANT: BARR, PHILIP J
;      APPLICANT: KIEFER, MICHAEL C
;      TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PACE 4 AND
;      TITLE OF INVENTION: PACE 4.1 GENE AND POLYPEPTIDES IN CELLS
;      NUMBER OF SEQUENCES: 16
;      CORRESPONDENCE ADDRESS:
;          ADDRESSEE: COOLEY GODWARD CASTRO HUDDLESON & TATUM
;          STREET: FIVE PALO ALTO SQUARE
;          CITY: PALO ALTO
;          STATE: CALIFORNIA
;          COUNTRY: USA
;          ZIP: 94306
;      COMPUTER READABLE FORM:
;          MEDIUM TYPE: Floppy disk
;          COMPUTER: IBM PC compatible
;          OPERATING SYSTEM: PC-DOS/MS-DOS
;          SOFTWARE: PatentIn Release #1.0, Version #1.25
;      CURRENT APPLICATION DATA:
;          APPLICATION NUMBER: PCT/US93/02147A
;          FILING DATE: 19930309
;          CLASSIFICATION:
;      PRIOR APPLICATION DATA:
;          APPLICATION NUMBER: US 07/848,629
;          FILING DATE: 09-MAR-1992
;      ATTORNEY/AGENT INFORMATION:
;          NAME: NEELEY PH.D., RICHARD L
;          REGISTRATION NUMBER: 30092
;          REFERENCE/DOCKET NUMBER: CHIR-009/00US
;      TELECOMMUNICATION INFORMATION:
;          TELEPHONE: (415) 494-7622
;          TELEFAX: (415) 857-0663
;          TELEX: 380816 COOLEY PA
;  INFORMATION FOR SEQ ID NO: 2:
;      SEQUENCE CHARACTERISTICS:
;          LENGTH: 969 amino acids
;          TYPE: amino acid
;          TOPOLOGY: linear
;      MOLECULE TYPE: protein
;  PCT-US93-02147A-2

Query Match     6.6%;   Score 166.5;  DB 5;   Length 969;
Best Local Similarity  20.7%;  Pred. No. 2.1e-06;
Matches   86;  Conservative   53;  Mismatches 135;   Indels 141;   Gaps  25;

Qy   70  PQKKDAGAQPNPPATANLV--TQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENA  127
              |   |  | :||:: :|       ||::|| ::|        |:|:||  |:|
Db  672  PEDEEDYTAQSTPGSANILQTSVCHPEC--GDKGCDGPNAD------QCLNC-VHF---SL  720

Qy  128  PNFNAGASTCTACPVNRVGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECV  187
              ||  |  | ::|      |  |  || |    |::
Db  721  GSVKTSRKCVSVCPLGYFG----DTAARRCRRCHKGCETCSS-------RAATQCL  765
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Query Match      6.2%;    Score 157;  DB 2;  Length 3111;
Best Local Similarity  18.5%;  Pred. No. 6.8e-05;
Matches  98;  Conservative  47;  Mismatches  164;  Indels  222;  Gaps  28;

Qy   18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA 77
         ::  |||:|  :|:|:     |                 |       :||
Db  721 VEVCQCPPGY--------------TGSSCESCWPRHRRVNGTIF---GGICEPC------- 757

Qy   78 QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRIN------------FYNE 125
         |      |         ||:|||:  |                 || ||||:
Db  758 ------------QC----------FGHAESCDDVTGECLNCKDHTGGPYCDKCLPGFYGE 795

Qy  126 NAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTE 185
         | |||:         ::|:  || :| |    :|||
Db  796 PTKGTSEDCQPC-ACPLN-----IPSNNFS--PTCHLDRSLG-LICDGCPVGY--TGPR 843

Qy  186 CVKCRLNFYYNGNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCN------V 239
         | ||:   :  ||           |  ||                         :|
Db  844 CERCAEGYF------GQPSVPGGS-CQPC--------QCNDNLDFSIPGSCDSLSGSCL 887

Qy  240 ACPDGTISAAGVNNWVAQNTECTNCAPNFYNNAPNFNFNPGNSTCLPCPANKDYGAEATAG 299
         ||  |     |             ||                       |  ||
Db  888 ICKPGT----------TGRYCELCADGYFGDAVDAKN----CQPCRCN--------AG 923

Qy  300 G---------------AATLAKQCNIACPDGTAIASGATNYV---------------- 326
         |               |||  |||   ||  || ||
Db  924 GSFSEVCHSQTGQCECRANVQGQRCD-KCKAGTFGLQSARGCVPCNCNSFGSKSFDCEES 982

Qy  327 --------ILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQC 378
                 |:|       ||          |:|       |||             :|
Db  983 GQCWCQPGVTGKKCDRCAHGYF-----NFQEGG--CTACECSHLGNNCDPKTG------RC 1030

Qy  379 ALECPAGTVLTDGTTSTYKQAASECVKCAANFY----------------TTKQTDWVAGIDT 424
         | |                 :|:  ||||                     :|
Db 1031 I--CPPNTI------------GERCSKCAPNTWGHSITTGCKACNCSTVGSLDFQCNVNT 1076

Qy  425 ------------CTSCNKK----------------LTSGAEANLPESAKKNIQC 450
                     ||:||:                 |:|
Db 1077 GQCNCHPKFSGAKCTECSRGHWNYPRCNLCDCFLPGTDATTCDSETKKCSC 1127
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RESULT 11
US-08-125-077-4
; Sequence 4, Application US/08125077
; Patent No. 5872231 5840863
; Patent No. 5872231 5840863
GENERAL INFORMATION:
    APPLICANT: Engvall, Eva
    APPLICANT: Leivo, Ilmo
    TITLE OF INVENTION: Nucleic Acids Encoding Merosin, Merosin
    TITLE OF INVENTION: Fragments and Uses Thereof
    NUMBER OF SEQUENCES: 23
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Campbell and Flores
    STREET: 4370 La Jolla Village Drive, Suite 700
    CITY: San Diego
    STATE: California
    COUNTRY: USA
    ZIP: 92122
    COMPUTER READABLE FORM:
    MEDIUM TYPE: Floppy disk
    COMPUTER: IBM PC compatible
    OPERATING SYSTEM: PC-DOS/MS-DOS
    SOFTWARE: PatentIn Release #1.0, Version #1.25
    CURRENT APPLICATION DATA:
    APPLICATION NUMBER: US/08/125,077
    FILING DATE: 22-SEP-1993
    CLASSIFICATION: 435
    PRIOR APPLICATION DATA:
    APPLICATION NUMBER: US PCT/US 94/10730
    FILING DATE: 21-SEP-1994
    PRIOR APPLICATION DATA:
    APPLICATION NUMBER: US 07/472,319

    FILING DATE: 30-JAN-1990
    PRIOR APPLICATION DATA:
    APPLICATION NUMBER: US 07/919,951
    FILING DATE: 27-JUL-1992
    ATTORNEY/AGENT INFORMATION:
    NAME: Campbell, Cathryn A.
    REGISTRATION NUMBER: 31,815
    REFERENCE/DOCKET NUMBER: P-LA 9721
    TELECOMUNICATION INFORMATION:
    TELEPHONE: (619) 535-9001
    TELEFAX: (619) 535-8949
    INFORMATION FOR SEQ ID NO: 4:
    SEQUENCE CHARACTERISTICS:
    LENGTH: 3111 amino acids
    TYPE: amino acid
    TOPOLOGY: linear
US-08-125-077-4

RESULT 12
US-08-317-450B-15
; Sequence 15, Application US/08317450B
; Patent No. 5660982
GENERAL INFORMATION:
    APPLICANT: Tryggvason, Karl
    APPLICANT: Kaliunki, Pekka
    APPLICANT: Pyke, Charles
    TITLE OF INVENTION: Laminin Chains: Diagnostic and
    TITLE OF INVENTION: Therapeutic Use
    NUMBER OF SEQUENCES: 19
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: BANNER & ALLEGRETTI, LTD.

STREET: Ten South Wacker Drive
CITY: Chicago
STATE: Illinois
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/317,450B
 FILING DATE: 04-OCT-1994
 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
 NAME: Chao, Mark
 REGISTRATION NUMBER: 37,293
 REFERENCE/DOCKET NUMBER: 94,778
TELECOMMUNICATION INFORMATION:
 TELEPHONE: 312-715-1000
 TELEFAX: 312-715-1234
INFORMATION FOR SEQ ID NO: 15:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 1111 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
US-08-317-450B-15

Query Match   5.9%; Score 150.5; DB 1; Length 1111;
Best Local Similarity 20.0%; Pred. No. 6.5e-05;
Matches 82; Conservative 49; Mismatches 138; Indels 141; Gaps 25;

Qy   18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA 77
          :: ||||: | | ::: | |||||
Db 379 VEQCICPVGYK---GQF-------CQDCASG--YKRDSARLGPFGTCIPCNCQGGGAC 424

Qy   78 QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC 137
          :| ||:|||| |||:|| |||| | :|
Db 425 DPD-------TGDCYSGDENP----DIECADCPIGFYND--PH--DPRSC 459

Qy 138 TACPVNRVGGALTAGNAATIVAQ------CNVACPTGTALDDGVTTDYVRSFTECVKCRLN 192
         || || || | ||:| ||: || ||| ||
Db 460 KPCPCHN-------GFSCSVIPETEEVVCN-NCPPGV------TGARCELCADG 499

Qy 193 FYYNGNNGNTPFNPGKSQCTPCAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVN 252
         | | | | : ||| | ::: ||| | | |:| |
Db 500 YFGDPFGEHGPVRP----CQPCQC--NSNVDPSASGNCDRLTGRC-LKCIHNT--AGI- 548

Qy 253 NWVAQNTECTNCAPNFYNNN-APNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCNIA 311
          :: | || | | | | | | | || :: |::
Db 549 ------YCDQCKAGYFGDPLAP--NPADK-CRACNCN-PMGSEP---------VG 584

Qy 312 C-PDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKAC------------P 357
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Db 585 CRSDGT----------------------------CVCKPGFGGPNCEHGAFSCPACYNQVKIQMDQFMQQ 626

Qy 358 ANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCA 407
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Db 627 LQRMEALISKAQGGDGVVPDTELE-------GRMQQAEQALQDILRDA 667

RESULT 13
US-08-800-593-15
; Sequence 15, Application US/08800593
; Patent No. 6143505
; GENERAL INFORMATION:
;   APPLICANT: Tryggvason, Karl
;   APPLICANT: Kallunki, Pekka
;   APPLICANT: Pyke, Charles
;   TITLE OF INVENTION: Laminin Chains: Diagnostic and
;   TITLE OF INVENTION: Therapeutic Use

NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
 ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
 STREET: 300 South Wacker Drive
 CITY: Chicago
 STATE: Illinois
 COUNTRY: USA
 ZIP: 60606
COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/800,593
 FILING DATE: 18-FEB-1997
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/317,450
 FILING DATE: 04-OCT-1994
 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
 NAME: Chao, Mark
 REGISTRATION NUMBER: 37,293
 REFERENCE/DOCKET NUMBER: 94,778-B
TELECOMMUNICATION INFORMATION:
 TELEPHONE: 312-913-0001
 TELEFAX: 312-913-0002
INFORMATION FOR SEQ ID NO: 15:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 1111 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
US-08-800-593-15

Query Match   5.9%; Score 150.5; DB 4; Length 1111;
Best Local Similarity 20.0%; Pred. No. 6.5e-05;
Matches 82; Conservative 49; Mismatches 138; Indels 141; Gaps 25;

Qy   18 IKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA 77
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Qy   78 QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC 137
          :| ||:|||| |||:|| |||| | :|
Db 425 DPD-------TGDCYSGDENP----DIECADCPIGFYND--PH--DPRSC 459

Qy 138 TACPVNRVGGALTAGNAATIVAQ------CNVACPTGTALDDGVTTDYVRSFTECVKCRLN 192
         || || || | ||:| ||: || ||| ||
Db 460 KPCPCHN-------GFSCSVIPETEEVVCN-NCPPGV------TGARCELCADG 499

Qy 193 FYYNGNNGNTPFNPGKSQCTPCAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVN 252
         | | | | : ||| | ::: ||| | | |:| |
Db 500 YFGDPFGEHGPVRP----CQPCQC--NSNVDPSASGNCDRLTGRC-LKCIHNT--AGI- 548

Qy 253 NWVAQNTECTNCAPNFYNNN-APNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCNIA 311
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Db 549 ------YCDQCKAGYFGDPLAP--NPADK-CRACNCN-PMGSEP---------VG 584

Qy 312 C-PDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKAC------------P 357
          | || | : | | ||| |: | | | || || || :::
Db 585 CRSDGT----------------------------CVCKPGFGGPNCEHGAFSCPACYNQVKIQMDQFMQQ 626

Qy 358 ANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCA 407
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Db 627 LQRMEALISKAQGGDGVVPDTELE-------GRMQQAEQALQDILRDA 667

RESULT 14
US-08-317-450B-13
; Sequence 13, Application US/08317450B

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Patent No. 5660982
; GENERAL INFORMATION:
;   APPLICANT: Tryggvason, Karl
;   APPLICANT: Kallunki, Pekka
;   APPLICANT: Pyke, Charles
;   TITLE OF INVENTION: Laminin Chains: Diagnostic and
;   TITLE OF INVENTION: Therapeutic Use
;   NUMBER OF SEQUENCES: 19
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: BANNER & ALLEGRETTI, LTD.
;     STREET: Ten South Wacker Drive
;     CITY: Chicago
;     STATE: Illinois
;     COUNTRY: USA
;     ZIP: 60606
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/317,450B
;   FILING DATE: 04-OCT-1994
;   CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
;   NAME: Chao, Mark
;   REGISTRATION NUMBER: 37,293
;   REFERENCE/DOCKET NUMBER: 94,778
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: 312-715-1000
;   TELEFAX: 312-715-1234
; INFORMATION FOR SEQ ID NO: 13:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 1193 amino acids
;     TYPE: amino acid
;     TOPOLOGY: linear
;     MOLECULE TYPE: protein
US-08-317-450B-13

Query Match            5.9%;  Score 150.5;  DB 1;  Length 1193;
Best Local Similarity  20.0%;  Pred. No. 7.1e-05;
Matches  82;  Conservative  49;  Mismatches 138;  Indels 141;  Gaps  25;

Qy    18  IKSANCPVGTETNTAGGVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGA  77
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Qy    78  QPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTC  137
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Db   425  DPD------------TGDCYSGDENP---DIECADCPIGFYND--PH--DPRSC  459

Qy   138  TACPVNRVGGALTAGNAATIVAQ----CNVACPTGTALDDGVTTDYVRSFTECVKCRLN  192
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Db   460  KPCPCHN------GFSCSVIPETEEVVCN-NCPPGV----------TGARCELCADG  499

Qy   193  FYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVN  252
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Qy   253  NWVAQNTECTNCAPNFYNNN-APNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCNIA  311
          |     |  |   | :| ||  ||  |  |  ||    || ||    |       :|:
Db   549  -----YCDQCKAGYFGDPLAP--NPADK-CRACNCN-PMGSEP---------VG  584

Qy   312  C-PDGTAIASGATNVILQTECLNCAANFYFDGNNFQAGSSRCKAC---------P  357
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Db   585  CRSDGT--------------CVCKPGFGGPNCEHGAFSCPACYNQVKIQMDQFMQQ  626

Qy   358  ANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCA  407
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Db   627  LQRMEALISKAQGGDGVVPDTELE-------GRMQQAEQALQDILRDA  667
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Wed Dec 26 12:01:32 2001

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Db    585  CRSDGT---------CVCKPGFGGPNCEHGAFSCPACYNQVKIQMDQFMQQ  626

Qy    358  ANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCA  407
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Db    627  LQRMEALISKAQGGDGVVPDTELE--------GRMQQAEQALQDILRDA   667


Search completed: December 26, 2001, 10:35:54
Job time: 145 sec
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